


# Repetitive Extragenic Palindromic (rep-PCR)

Highly-conserved repeated DNA elements

~ 500 copies of the REP consensus sequence in the  
E. coli chromosome.

# rep-PCR

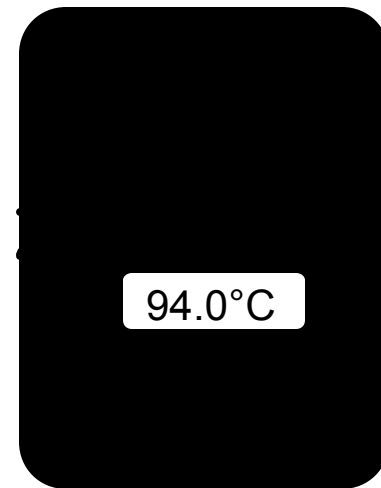
#1 



#2 



#3 

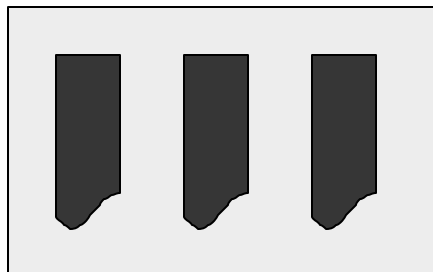
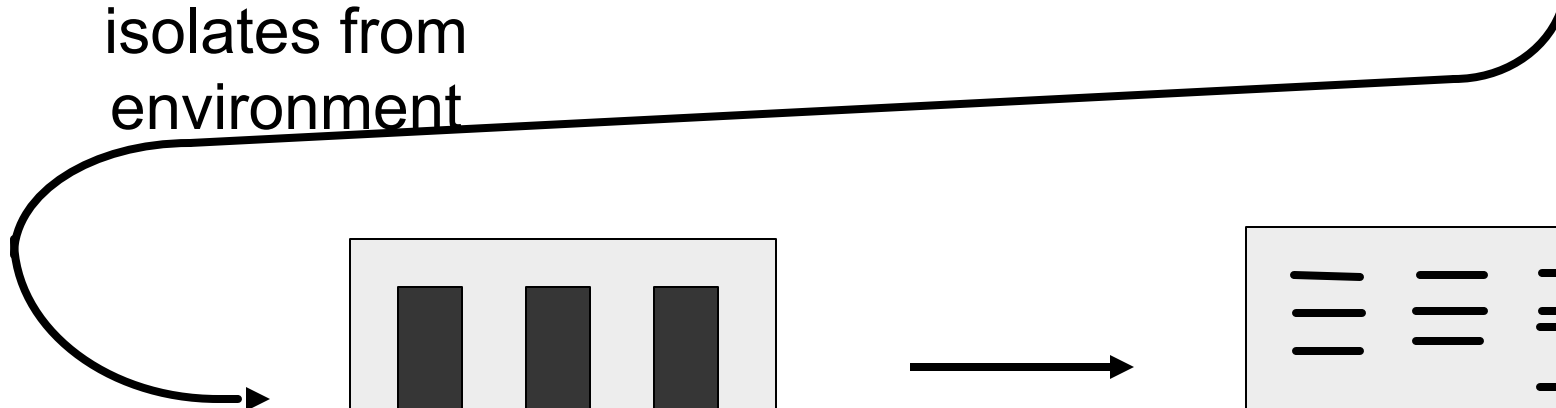


94.0°C

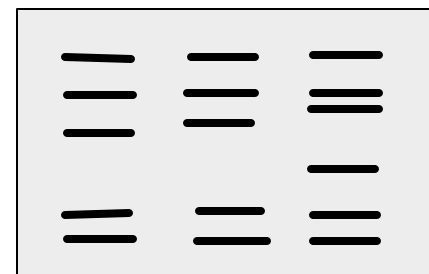
Three *E. coli*  
isolates from  
environment

Lyse cells

Amplify DNA

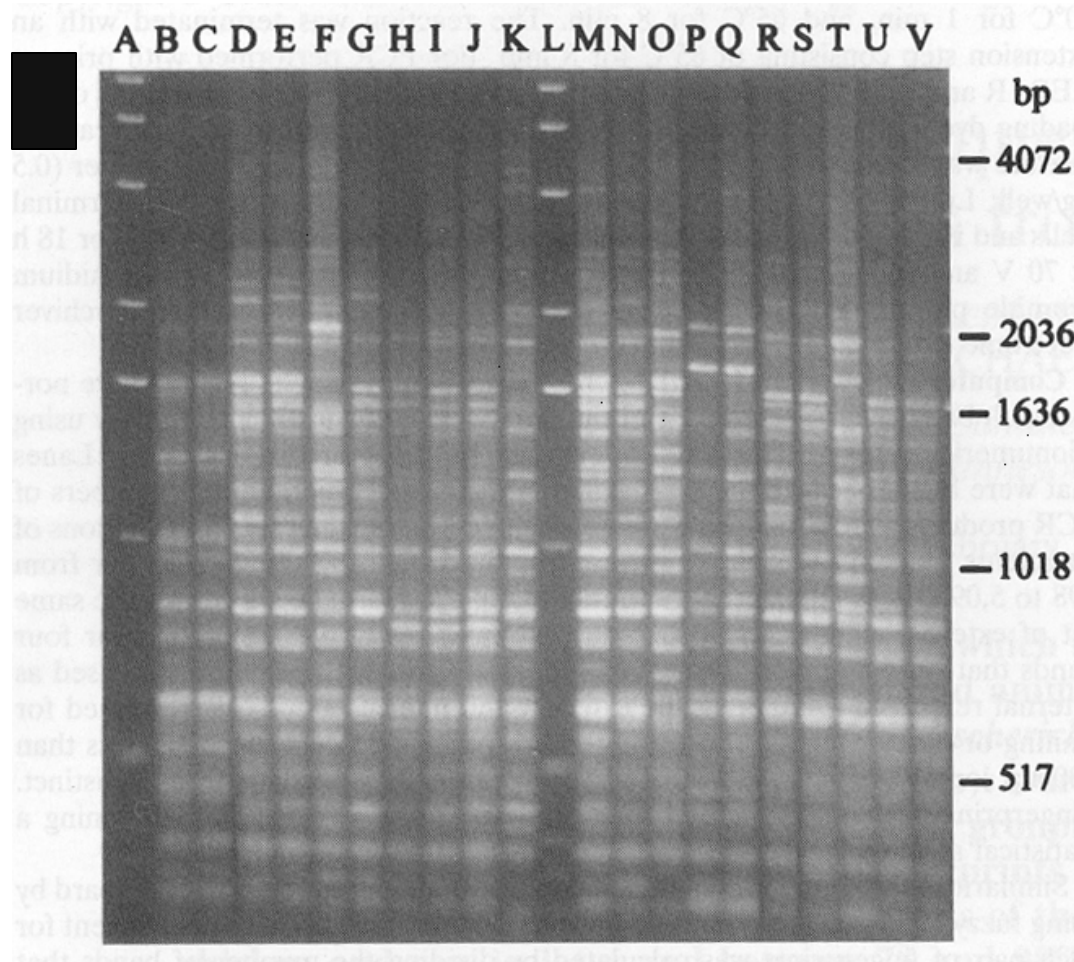


Run DNA on gel



Stain DNA to give  
“fingerprint”

# rep-PCR DNA Fingerprint



Adapted from Dombek et al. (2000)

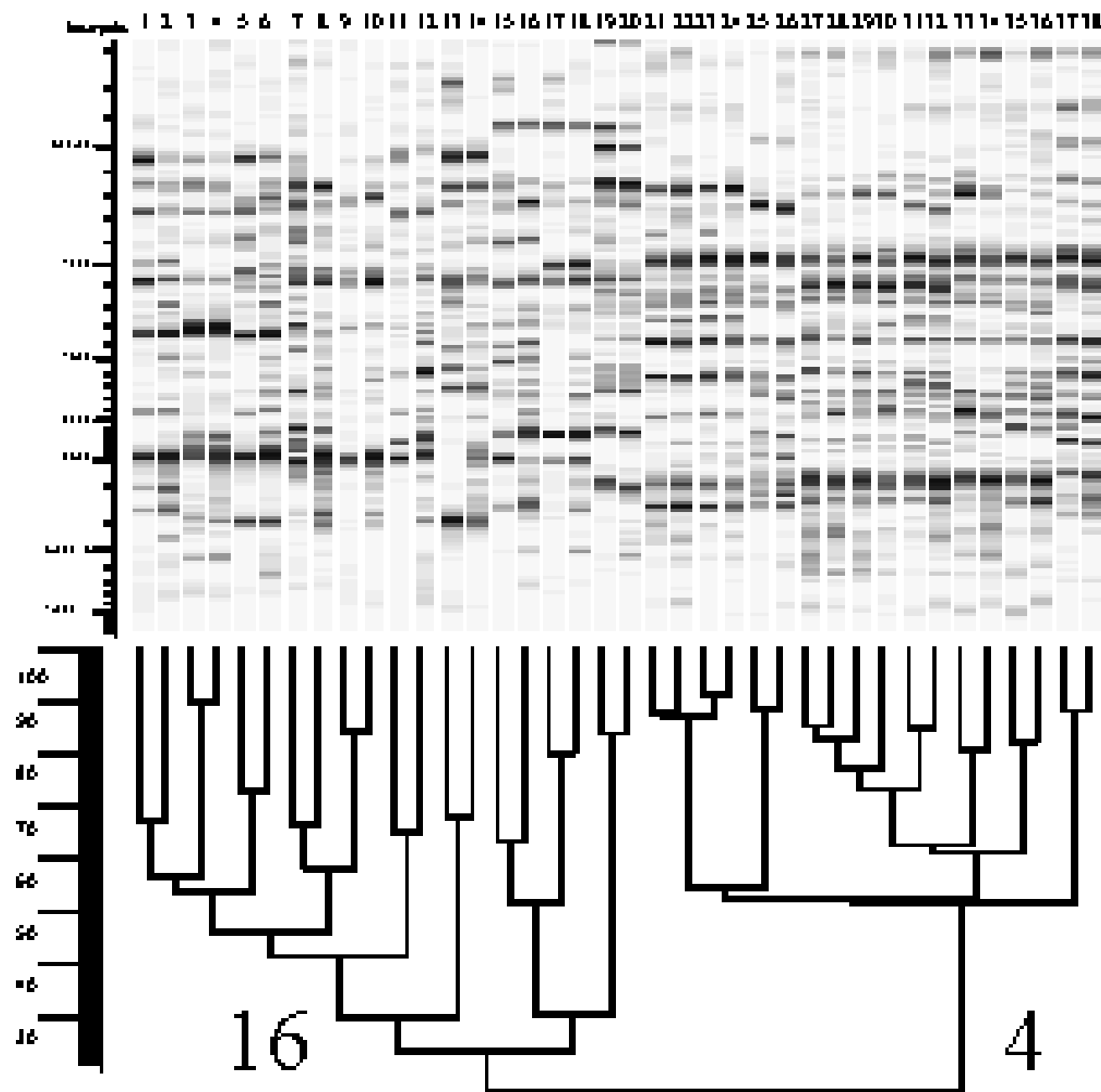


Fig. 7. Cluster analysis of REP-PCR genomic fingerprints of 38 (non specified) *Xanthomonas* strains belonging to DNA homology groups 4 and 16 (Vauterin *et al.* 1995) using the product moment and UPGMA method.

# rep-PCR

## Advantages

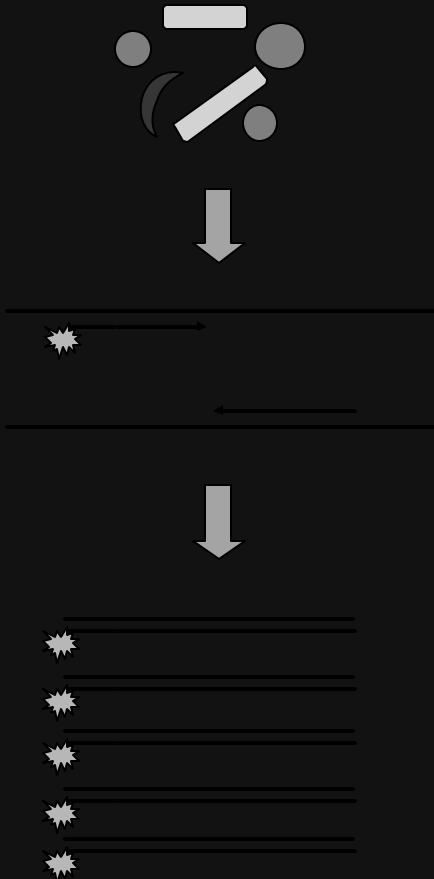
- Highly reproducible
- Excellent discrimination

## Disadvantages

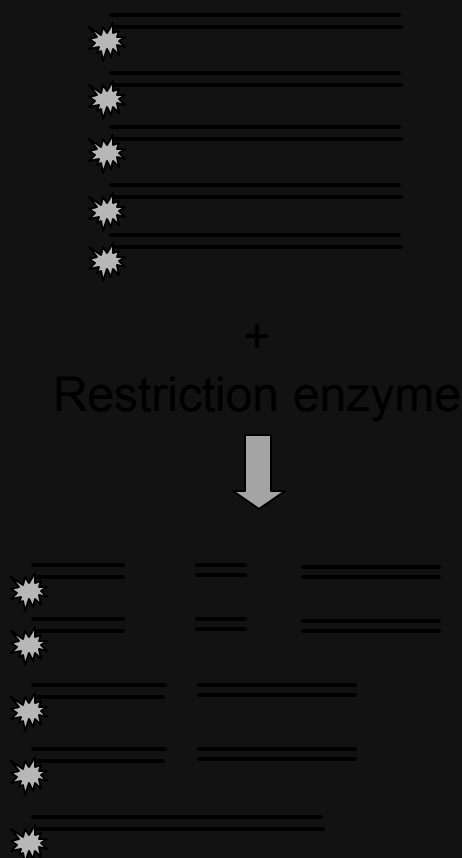
- Knowledge of DNA sequence needed
- Amplifying errors

# T-RFLP (terminal restriction fragment polymorphism)

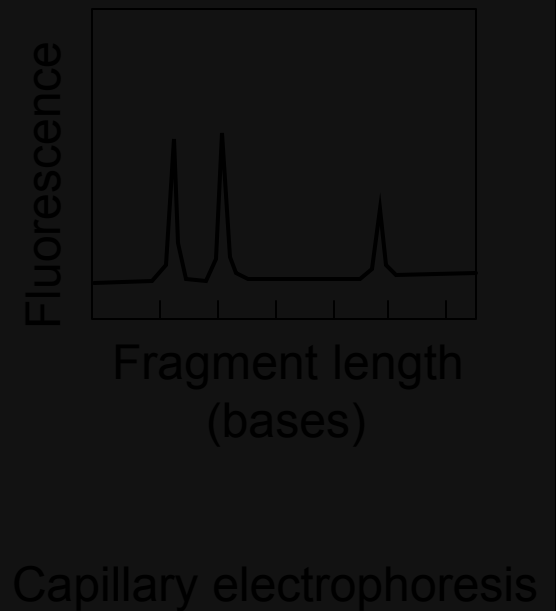
1) PCR-amplify and label target gene(s):



2) Digest labeled PCR products



3) Detect labeled terminal fragments



PCR = polymerase chain reaction

# Terminal Restriction Fragment Length Polymorphism (T-RFLP)

## Advantages

Does not require  
culturing

No library needed

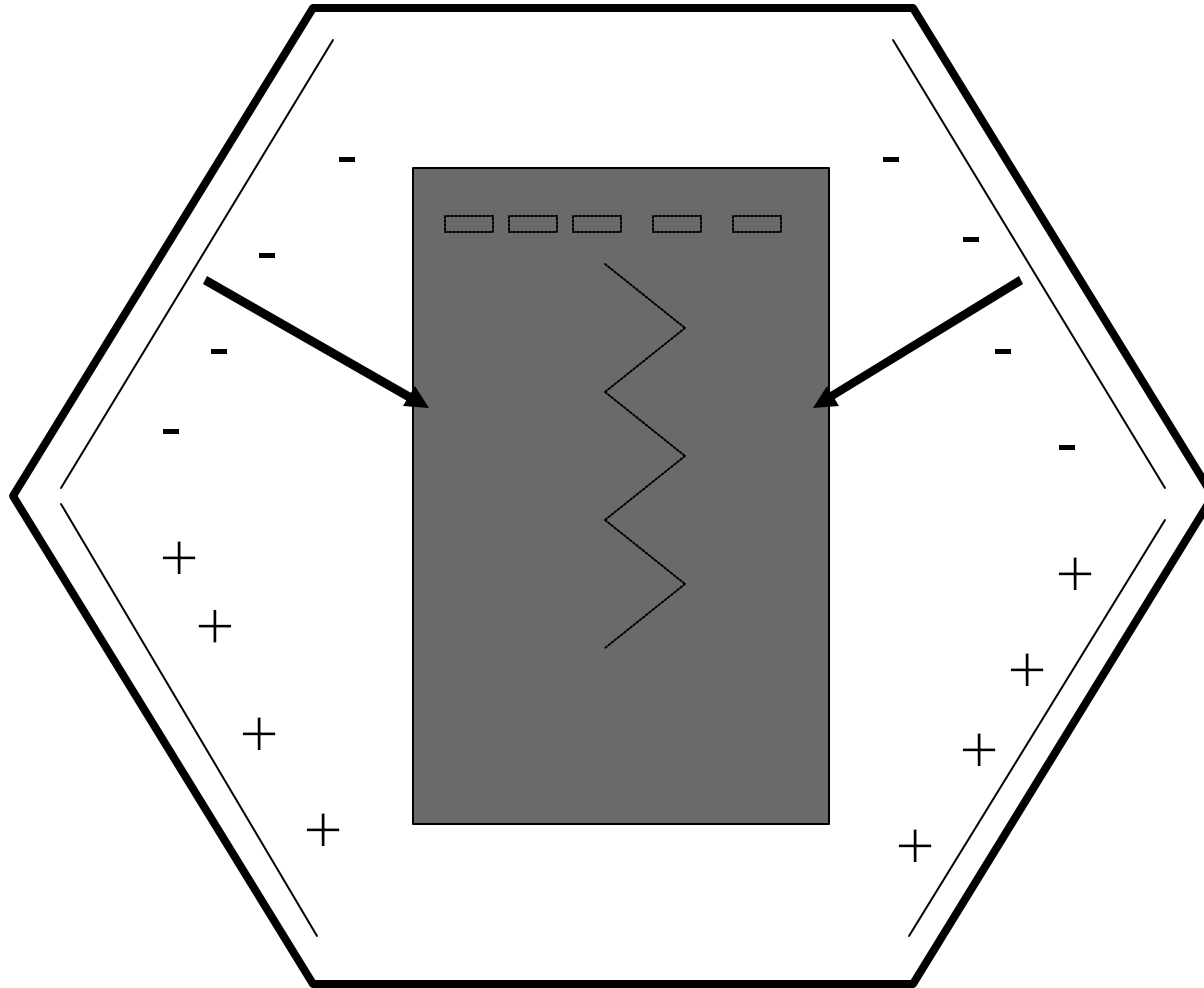
## Disadvantages

Expensive  
equipment needed

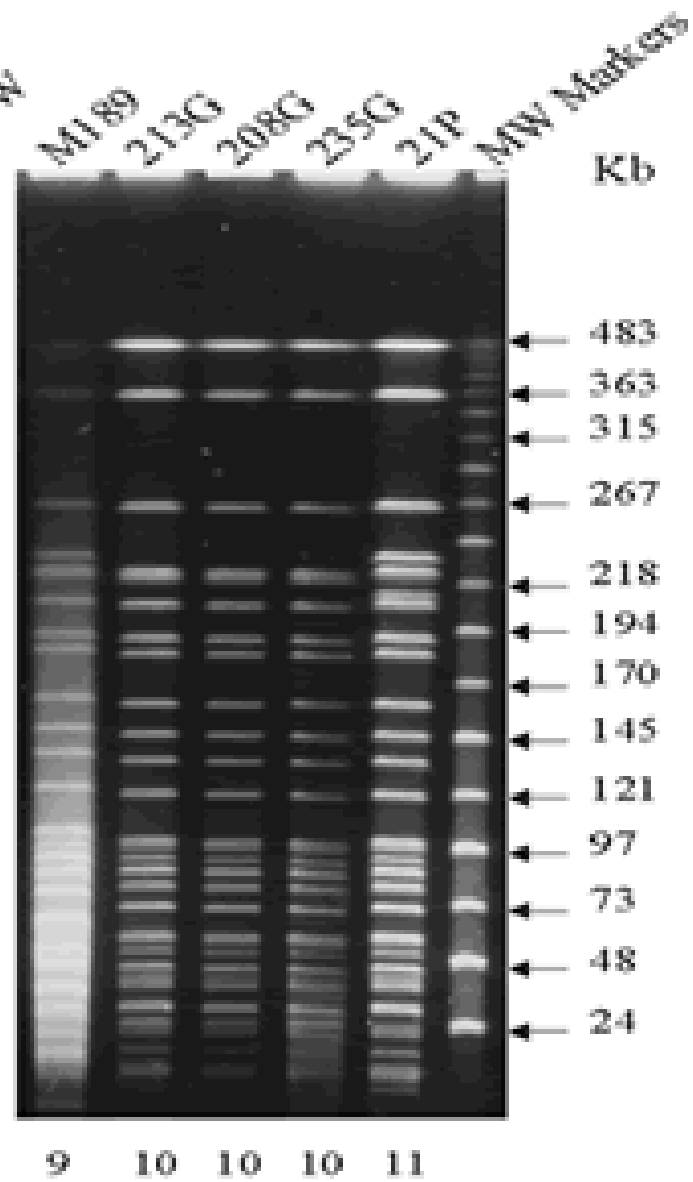
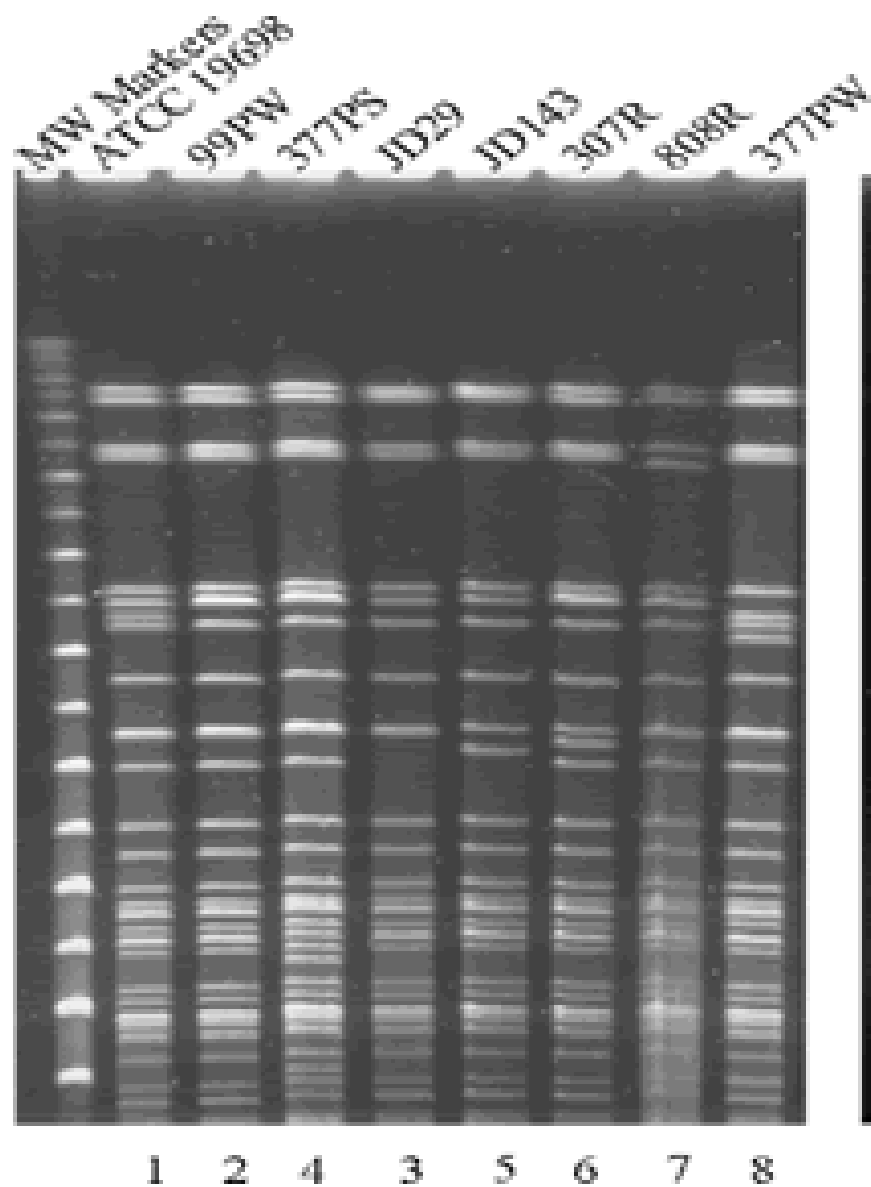
Technically  
demanding

# Pulse Field Gel Electrophoresis

Electric Field 1      Switch Time      Electric Field 2







# Pulse Field Electrophoresis (PFGE)

## Advantages

Used in genotyping  
and epidemiology

High discrimination

Reproducible

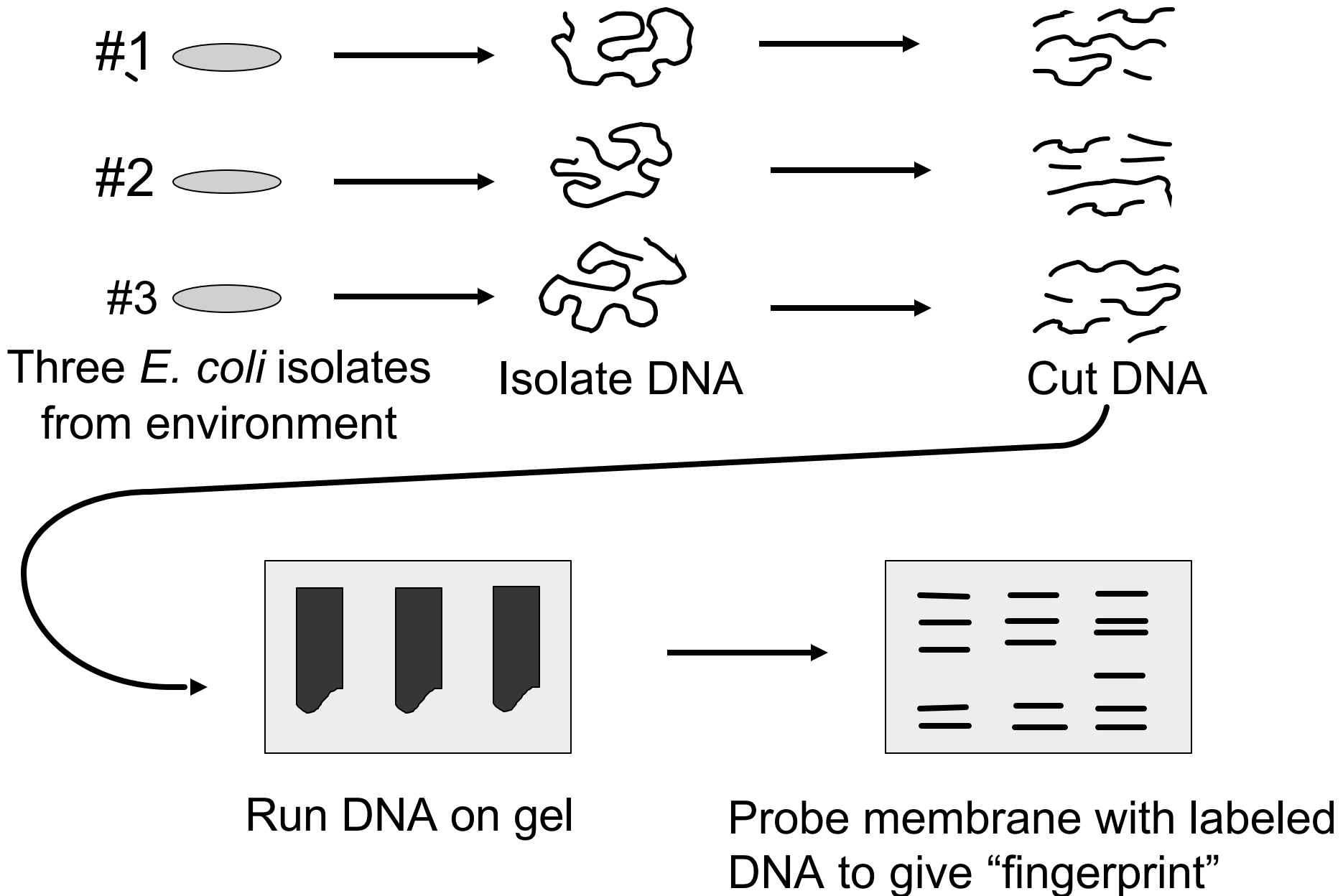
Conclusive results

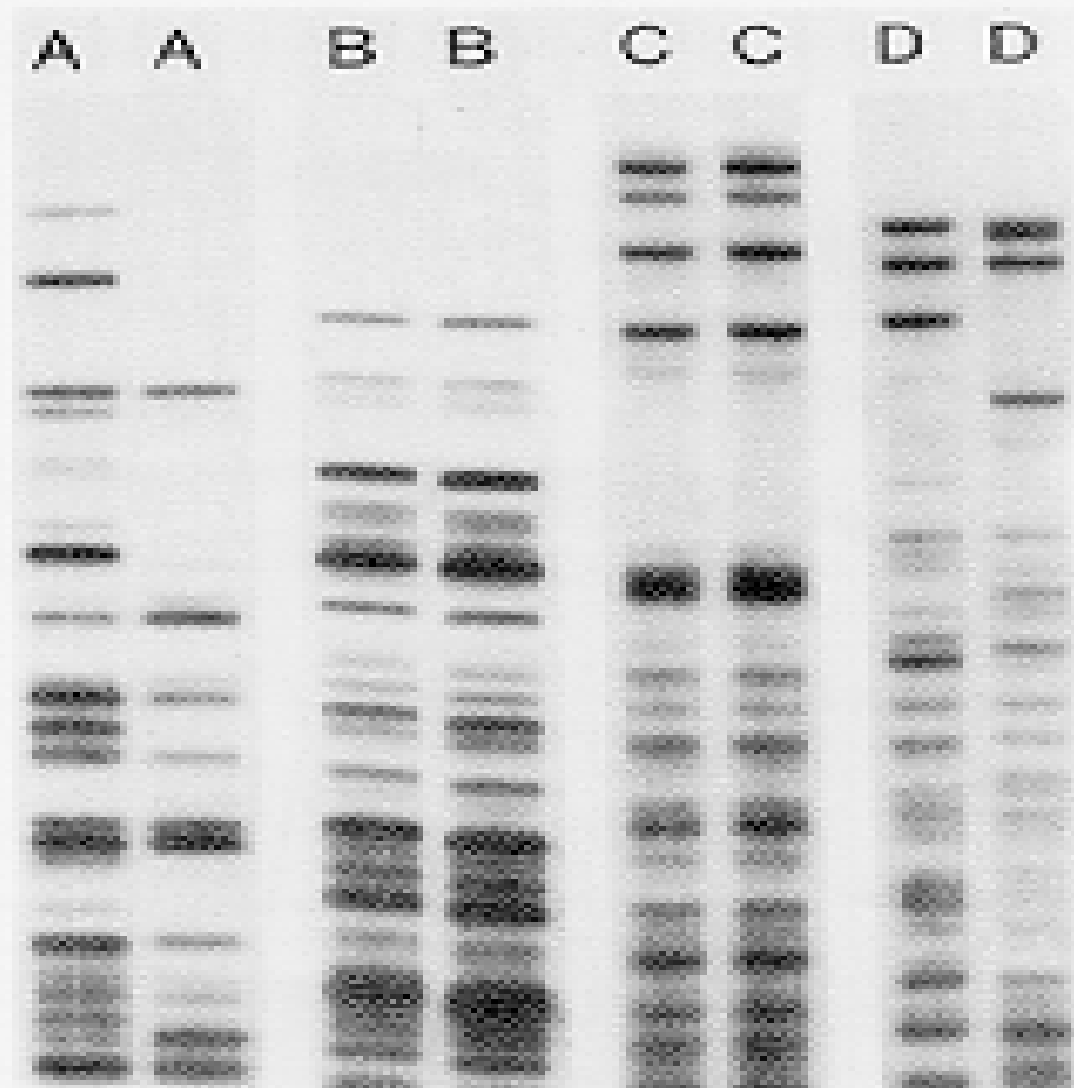
## Disadvantages

Long assay time

Limited Simultaneous  
strains processing

# Ribotyping





# Ribotyping

## Advantages

- Easy to type
- Highly reproducible
- Easy to perform
- Easy to interpret
- Easy to automate

## Disadvantages

- Tedious
- Time-consuming
- Expensive

# MST Methods

## Perception vs. Reality

- Great interest and need for a reliable molecular method for MST.
- Many methods being studied but none fulfills the need very well. Not ready for primetime!
- All potential MST molecular methods require well trained analysts, are technically challenging and are expensive.
- MST research needs are nearly overwhelming.

# MST Methods Research Needs

- Host origin database
- Biogeographic variability
- Temporal variability
- Environmental factors (e.g., rainfall)
- Indicator organisms
- Automation